## LIST OF SEQUENCES

5 MA3 / GENERAL INFORMATION:

APPLICANT:

NAMA: INSTITUTO CIENTIFICO Y TECNOLOGICO

DE NAVARRA, S.A.

STREET\ Avenida Pío XII, 53

TOWN: Pamplona

STATE OR PROVINCE: Navarra

COUNTRY: Spain

POST CODE: \31008

TELEPHONE: 948-10 56 00

FAX: 948-17 52\23

TITLE: "USE OF ALPHA\5 INTERFERON IN THE TREATMENT OF VIRAL

LIVER DISEASES".

NUMBER OF SEQUENCES: 1

ADDRESS FOR CORRESPONDENCE

NAME: INSTITUTO CIENTIFICO Y TECNOLOGICO

DE NAVARRA, S.A.

STREET: Avenida Pío XII, 53

TOWN: Pamplona

STATE OR PROVINCE: Navarra

COUNTRY: Spain
POST CODE: 31008

COMPUTER READABLE FORM:

TYPE OF MEDIUM: 3.5" DISK

COMPUTER: PC

OPERATING SYSTEM: WINDOWS

WORD PROCESSOR: WORD

INFORMATION ON THE LAWYER/AGENT:

NAME: ALBERTO DE ELZABURU

REGISTER NUMBER: 232/1

REFERENCE REGISTER NUMBER: P-99043

TELECOMMUNICATIONS INFORMATION:

TELEPHONE: 91 7009400

FAX: 91 3193810

TELEX OR ELECTRONIC MAIL: elzaburu@elzaburu.es

INFORMATION FOR SEQ ID NO.: 1

CHARACTERISTICS OF THE SEQUENCE:

\_LENGTH:\_274 base\_pairs -

TYPE: nucleotides

NUMBER OF STRANDS:

CONFIGURATION: linear

TYPE OF MOLECULE: cDNA

HYPOTHETICAL: NO

ANTI-DIRECTION: NO

SOURCE OF ORIGIN:

Organism: Homo Sapiens

Type of tissue: liver

POSITION IN THE GENOME: Chromosome 9

CHARACTERISTIC:

NAME/KEY: IFN $\alpha$ 5

Method of identification: RT-PCR, Sequencing

OTHER INFORMATION:

Nucleotides 672 to 945 in the sequence of the IFN $\alpha$ 5 gene published in the Genbank database under access number X02956.

DESCRIPTION OF THE SEQUENCE SEQ ID NO: 1

TC CAT GAG ATG	ATC CAG CAG ACC	TTC AAT CTC TTC	AGC ACA AAG GAC TCA	50
His Glu Met	Ile Gln Gln Thr	r Phe Asn Leu Phe	Ser Thr Lys Asp Ser	
1	5	10	15	
TCT GCT ACT TGG	G GAT GAG ACA CT	T CTA GAC AAA TTC	TAC ACT GAA CTT TAC	101
Ser Ala Thr Trp	Asp Glu. Thr Le	u Leu Asp Lys Phe	Tyr Thr Glu Leu Tyr	
20	\	25	30	
CAG CAG CTG AAT	GAC CTG GAA GC	CTGT ATG ATG CAG	GAG GTT GGA GTG GAA	152
Gln Gln Leu Asn	Asp Leu Glu Ala	a Cys Met Met Gln	Glu Val Gly Val Glu	
35	40	45	50	
GAC ACT CCT CTG	ATG AAT GTG GA	ATC CTG ACT	GTG AGA AAA TAC TTT	203
Asp Thr Pro Leu	ı Met Asn Val Ası	sp Ser Ite Leu Thr	Val Arg Lys Tyr Phe	
	5.5			
CAA AGA ATC ACC	C CTC TAT CTG AC	CA GAG AAG AAA TAC	AGC CCT TGT GCA TGG	254
Gln Arg Ile Thr	Leu Tyr Leu Th	ır Glu Lys Dys Tyr	Ser Pro Cys Ala Trp	
70	7.	75	80	
GAG GTT GTC AGA	A GCA GAA AT	. \		274
Glu Val Val Arg	, Ala Glu	\		
85	90			